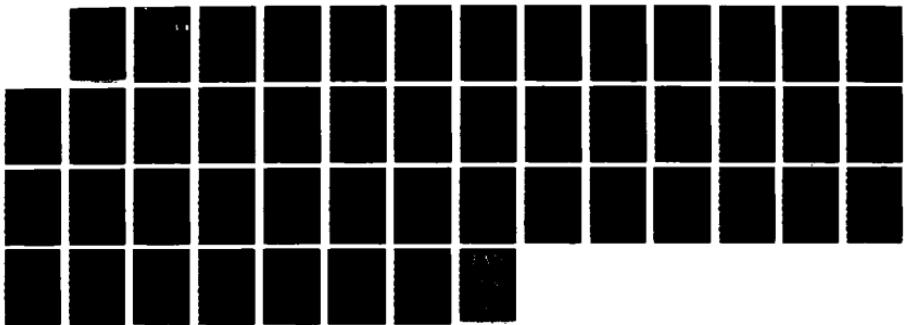
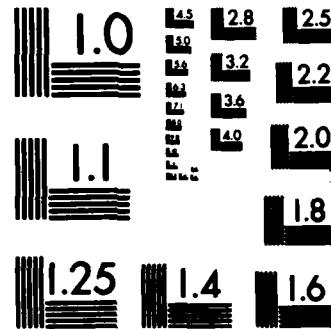


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ANNOTATED COMPUTER OUTPUT FOR ILLUSTRATIVE EXAMPLES
OF CLUSTERING USING THE MIXTURE METHOD AND
TWO COMPARABLE METHODS FROM SAS

K.E. BASFORD, N.J. MILES-MCDERMOTT, AND W.T. FEDERER

February 1987

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**ANNOTATED COMPUTER OUTPUT FOR ILLUSTRATIVE EXAMPLES
OF CLUSTERING USING THE MIXTURE METHOD AND
TWO COMPARABLE METHODS FROM SAS**

K.E. BASFORD, N.J. MILES-MCDERMOTT, AND W.T. FEDERER

February 1987



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ABSTRACT

This is the annotated computer output for the three clustering methods considered in the associated technical report, BU-921-M and '87-5, Illustrative Examples of Clustering using the Mixture Method and Two Comparable Methods from SAS, by K.E. Basford, W.T. Federer, and N.J. Miles-McDermott. The computer output for the normal mixture model method is generated from a fortran program, KMM, written by K.E. Basford. Two other clustering methods are considered and are from SAS/CLUSTER, Version 5. These are Ward's method and the EML method. Two real data sets are processed.

COMMENTS

The annotated output should be read in sequence because explanations made on earlier pages are not necessarily repeated subsequently. Some pages may be composites of more than one output page, and some output pages are omitted because they are generally not useful for the purpose at hand. A general description of the mixture model approach to clustering is explained in detail and discussed in relation to other clustering methods in Basford (1986). SAS program documentation is in SAS User's Guide (1985a and b). Program documentation for KMM is available from K.E.

Basford and will appear in a forthcoming book by McLachlan and Basford (1987).

The data are presented below. Following on pages 9-10 the KMM and SAS control language for each example is presented. Control language is given in capital type with accompanying descriptions and notes given in boldface type. Program output follows on pages 12-44 with annotations in boldface and lower case type that describe output values in some detail.

DATA SETS

Two data sets are used for each of the three clustering methods presented. The first data set was taken from Habbema, Hermans, and van den Broek (1974). These examples are labeled CL-1-Habbema through CL-3-Habbema on the output pages. The second data set is the well known Iris data published by Fisher (1936). These examples are labeled CL-1-Fisher through CL-3-Fisher on the output pages. For each data set, the first example CL-1- illustrates the normal mixture model method of clustering using the KMM program. CL-2- illustrates Ward's method using SAS and CL-3- illustrates the EML method also using SAS.

The data taken from Habbema et al., consists of 45 observations on known haemophilia A carriers and 35

observations on known noncarriers. These data are shown in Table 1 and contain three variables. GROUP indicates whether the individual is a carrier (coded 2) or noncarrier (coded 1). The two other variables are used to discriminate between the normal individuals and the carriers in the clustering programs and are \log_{10} (AHF activity) and \log_{10} (AHF-like antigen). These variables were named ACTIVITY and ANTIGEN, respectively.

TABLE 1: Habbema et al., Haemophilia Data

GROUP	ACTIVITY	ANTIGEN
1	-0.00559	-0.16571
1	-0.16980	-0.15852
1	-0.34689	-0.18791
1	-0.08944	0.00642
1	-0.16791	0.07129
1	-0.08362	0.01059
1	-0.19789	-0.00054
1	-0.07621	0.03919
1	-0.19129	-0.21229
1	-0.10919	-0.11904
1	-0.52677	-0.47734
1	-0.08419	0.02482
1	-0.02252	-0.05805
1	0.00841	0.07821
1	-0.18266	-0.11384
1	0.12366	0.21397
1	-0.47022	-0.30989
1	-0.15191	-0.06864
1	0.00061	-0.11531
1	-0.20154	-0.04976
1	-0.19318	-0.22933
1	0.15069	0.09331
1	-0.12591	-0.06686
1	-0.15508	-0.12321
1	-0.19515	-0.10067
1	0.02908	0.04419

1	-0.22282	-0.17099
1	-0.09971	-0.07333
1	-0.19724	-0.06074
1	-0.08670	-0.05597
2	-0.49859	-0.08602
2	-0.50145	-0.29844
2	-0.13259	0.00970
2	-0.34787	-0.17209
2	-0.37553	-0.18652
2	-0.24466	-0.04067
2	-0.22047	0.00455
2	-0.21539	-0.02191
2	-0.25404	-0.05729
2	-0.37780	-0.26816
2	-0.06391	0.15694
2	-0.33510	-0.13676
2	-0.01493	0.15392
2	-0.03124	0.14001
2	-0.17402	-0.07764
2	-0.09636	0.05307
2	-0.02344	0.08038
2	-0.40546	-0.24184
2	-0.34776	0.11506
2	-0.36180	-0.20082
2	-0.69112	-0.33899
2	-0.36083	0.12372
2	-0.45348	-0.16817
2	-0.35388	0.07219
2	-0.47186	-0.10786
2	-0.36097	-0.03994
2	-0.32261	0.16697
2	-0.43193	-0.06869
2	-0.27342	-0.00203
2	-0.55728	0.05480
2	-0.49503	-0.01529
2	-0.51066	-0.24825
2	-0.16516	0.21321
2	-0.42318	-0.09981
2	-0.23746	0.28763
2	-0.34470	0.00969
2	-0.40465	-0.11618
2	-0.14158	0.16416
2	-0.15082	0.11372
2	-0.26421	0.08669
2	-0.33525	0.08753
2	-0.18782	0.25096
2	-0.17443	0.18924
2	-0.24443	0.16137
2	-0.47837	0.02821

The Fisher Iris data is shown in Table 2 and consists of four measurements on 50 plants from each of three species

of Iris: *Iris setosa*, *Iris versicolor*, and *Iris virginica*. These species were coded 1, 2, and 3, respectively, with a variable name of GROUP. The four measurement variables input into the clustering programs were sepal length (SLENGTH), sepal width (SWIDTH), petal length (PLENGTH), and petal width (PWIDTH).

TABLE 2: Fisher Iris Data

GROUP	SLENGTH	SWIDTH	PLENGTH	PWIDTH
1	5.1	3.5	1.4	0.3
1	4.4	3.2	1.3	0.2
1	4.4	3.0	1.3	0.2
1	5.0	3.5	1.6	0.6
1	5.1	3.8	1.6	0.2
1	4.9	3.1	1.5	0.2
1	5.0	3.2	1.2	0.2
1	4.6	3.2	1.4	0.2
1	5.0	3.3	1.4	0.2
1	4.8	3.4	1.9	0.2
1	4.8	3.0	1.4	0.1
1	5.0	3.5	1.3	0.3
1	5.1	3.3	1.7	0.5
1	5.0	3.4	1.5	0.2
1	5.1	3.8	1.9	0.4
1	4.9	3.0	1.4	0.2
1	5.3	3.7	1.5	0.2
1	4.3	3.0	1.1	0.1
1	5.5	3.5	1.3	0.2
1	4.8	3.4	1.6	0.2
1	5.2	3.4	1.4	0.2
1	4.8	3.1	1.6	0.2
1	4.9	3.6	1.4	0.1
1	4.6	3.1	1.5	0.2
1	5.7	4.4	1.5	0.4
1	5.7	3.8	1.7	0.3
1	4.8	3.0	1.4	0.3
1	5.2	4.1	1.5	0.1
1	4.7	3.2	1.6	0.2
1	4.5	2.3	1.3	0.3
1	5.4	3.4	1.7	0.2
1	5.0	3.0	1.6	0.2
1	4.6	3.4	1.4	0.3
1	5.4	3.9	1.3	0.4

1	5.0	3.6	1.4	0.2
1	5.4	3.9	1.7	0.4
1	4.6	3.6	1.0	0.2
1	5.1	3.8	1.5	0.3
1	5.8	4.0	1.2	0.2
1	5.4	3.7	1.5	0.2
1	5.0	3.4	1.6	0.4
1	5.4	3.4	1.5	0.4
1	5.1	3.7	1.5	0.4
1	4.4	2.9	1.4	0.2
1	5.5	4.2	1.4	0.2
1	5.1	3.4	1.5	0.2
1	4.7	3.2	1.3	0.2
1	4.9	3.1	1.5	0.1
1	5.2	3.5	1.5	0.2
1	5.1	3.5	1.4	0.2
2	6.4	3.2	4.5	1.5
2	5.5	2.4	3.8	1.1
2	5.7	2.9	4.2	1.3
2	5.7	3.0	4.2	1.2
2	5.6	2.9	3.6	1.3
2	7.0	3.2	4.7	1.4
2	6.8	2.8	4.8	1.4
2	6.1	2.8	4.7	1.2
2	4.9	2.4	3.3	1.0
2	5.8	2.7	3.9	1.2
2	5.8	2.6	4.0	1.2
2	5.5	2.4	3.7	1.0
2	6.7	3.0	5.0	1.7
2	5.7	2.8	4.1	1.3
2	6.7	3.1	4.4	1.4
2	5.5	2.3	4.0	1.3
2	5.1	2.5	3.0	1.1
2	6.6	2.9	4.6	1.3
2	5.0	2.3	3.3	1.0
2	6.9	3.1	4.9	1.5
2	5.0	2.0	3.5	1.0
2	5.6	3.0	4.5	1.5
2	5.6	3.0	4.1	1.3
2	5.8	2.7	4.1	1.0
2	6.3	2.3	4.4	1.3
2	6.1	3.0	4.6	1.4
2	5.9	3.0	4.2	1.5
2	6.0	2.7	5.1	1.6
2	5.6	2.5	3.9	1.1
2	6.7	3.1	4.7	1.5
2	6.2	2.2	4.5	1.5
2	5.9	3.2	4.8	1.8
2	6.3	2.5	4.9	1.5
2	6.0	2.9	4.5	1.5
2	5.6	2.7	4.2	1.3
2	6.2	2.9	4.3	1.3

2	6.0	3.4	4.5	1.6
2	6.5	2.8	4.6	1.5
2	5.7	2.8	4.5	1.3
2	6.1	2.9	4.7	1.4
2	5.5	2.5	4.0	1.3
2	5.5	2.6	4.4	1.2
2	5.4	3.0	4.5	1.5
2	6.3	3.3	4.7	1.6
2	5.2	2.7	3.9	1.4
2	6.4	2.9	4.3	1.3
2	6.6	3.0	4.4	1.4
2	5.7	2.6	3.5	1.0
2	6.1	2.8	4.0	1.3
2	6.0	2.2	4.0	1.0
3	6.3	3.3	6.0	2.5
3	6.7	3.3	5.7	2.1
3	7.2	3.6	6.1	2.5
3	7.7	3.8	6.7	2.2
3	7.2	3.0	5.8	1.6
3	7.4	2.8	6.1	1.9
3	7.6	3.0	6.6	2.1
3	7.7	2.8	6.7	2.0
3	6.2	3.4	5.4	2.3
3	7.7	3.0	6.1	2.3
3	6.8	3.0	5.5	2.1
3	6.4	2.7	5.3	1.9
3	5.7	2.5	5.0	2.0
3	6.9	3.1	5.1	2.3
3	5.9	3.0	5.1	1.8
3	6.3	3.4	5.6	2.4
3	5.8	2.7	5.1	1.9
3	6.3	2.7	4.9	1.8
3	6.0	3.0	4.8	1.8
3	7.2	3.2	6.0	1.8
3	6.2	2.8	4.8	1.8
3	6.9	3.1	5.4	2.1
3	6.7	3.1	5.6	2.4
3	6.4	3.1	5.5	1.8
3	5.8	2.7	5.1	1.9
3	6.1	3.0	4.9	1.8
3	6.0	2.2	5.0	1.5
3	6.4	3.2	5.3	2.3
3	5.8	2.8	5.1	2.4
3	6.9	3.2	5.7	2.3
3	6.7	3.0	5.2	2.3
3	7.7	2.6	6.9	2.3
3	6.3	2.8	5.1	1.5
3	6.5	3.0	5.2	2.0
3	7.9	3.8	6.4	2.0
3	6.1	2.6	5.6	1.4
3	6.4	2.8	5.6	2.1
3	6.3	2.5	5.0	1.9

3	4.9	2.5	4.5	1.7
3	6.8	3.2	5.9	2.3
3	7.1	3.0	5.9	2.1
3	6.7	3.3	5.7	2.5
3	6.3	2.9	5.6	1.8
3	6.5	3.0	5.5	1.8
3	6.5	3.0	5.8	2.2
3	7.3	2.9	6.3	1.8
3	6.7	2.5	5.8	1.8
3	5.6	2.8	4.9	2.0
3	6.4	2.8	5.6	2.2
3	6.5	3.2	5.1	2.0

Control Language

CL-1-Habbema (Mixture method from KMM)

```
75 2           ⇒ 75 is the number of observations and 2 is
                the number of variables
-0.005595 -0.165712
-0.169805 -0.158521      } INPUT DATA: ACTIVITY and ANTIGEN
:
:
-0.478366  0.028215
2           ⇒ number of clusters to be formed
2           ⇒ instructs KMM to assume unequal covariance matrices
1           ⇒ signals KMM that initial grouping estimates follow
1 1 2 1 1 1 1 1 1 1
2 1 1 1 1 1 2 1 1
1 1 1 1 1 1 1 1 1
2 2 1 2 2 1 1 1 1 2      } Initial groupings of observations
1 2 1 1 1 1 2 1 2      } (results of Ward's method were used)
2 1 1 1 2 1 1 2 1 2
2 1 2 2 1 1 2 1 1 1
1 2 1 1 2
```

CL-2-Habbema (Ward's method from SAS)

```
DATA GJ;
INPUT ACTIVITY ANTIGEN;      ⇒ Input variables
IF N LE 30 THEN GROUP=1;      } Defines the GROUP variable
ELSE GROUP=2;
CARDS;                         ⇒ Signals SAS that the data follow
-0.005595 -0.165712
-0.169805 -0.158521
:
:
-0.478366  0.028215
PROC CLUSTER OUTTREE=TREE METHOD=WARD;  ⇒ Requests CLUSTER analysis
                                         using Ward's method on ACTIVITY and ANTIGEN
VAR ACTIVITY ANTIGEN;
COPY GROUP;
PROC TREE SORT HEIGHT=N;      ⇒ Requests the Cluster Tree from 1 to n
                                         (75) clusters
ID GROUP;
PROC TREE NCL=2 OUT=OUT NOPRINT;
ID GROUP;                      } Causes SAS to produce 2x2
PROC FREQ;                     } table showing misclassifications
TABLE CLUSTER*GROUP;
```

CL-3-Habbema (EML method from SAS)

Same control language as for 2) above except substitute EML for WARD on PROC CLUSTER line.

CL-1-Fisher (Mixture method from KMM)

150 4 ⇒ 150 is number of observations and 4 is the number of variables
5.1 3.5 1.4 0.3
4.4 3.2 1.3 0.2 } Input data
:
6.5 3.2 5.1 2.0
3 ⇒ Number of clusters to be formed
1 ⇒ Instructs KMM to assume equal covariance matrices
1 ⇒ Signals KMM that initial grouping estimates follow
1 1 1 1 1 1 1 1 1 1 } Initial grouping of observations
1 1 1 1 1 1 1 1 1 1 } (results of Ward's method were used)
:
3 3 3 3 3 3 3 2 3 3

CL-2-Fisher (Ward's method from SAS)

DATA ONE;
INPUT SLENGTH SWIDTH PLENGTH PWIDTH; ⇒ Input variables
IF N LE 50 THEN GROUP=1;
ELSE IF N LE 100 THEN GROUP=2; } Defines the GROUP variable
ELSE GROUP=3;
CARDS; ⇒ Signals SAS that the data follow
5.1 3.5 1.4 0.3
4.4 3.2 1.3 0.2
:
6.5 3.2 5.1 2.0
PROC CLUSTER OUTTREE=TREE METHOD=WARD; } Requests the Cluster analysis
VAR SLENGTH SWIDTH PLENGTH PWIDTH; } using Ward's method on the 4
COPY GROUP; } variables SLENGTH, SWIDTH,
 } PLENGTH, AND PWIDTH
PROC TREE DATA=TREE SORT HEIGHT=N; } Requests cluster
ID GROUP; } tree
PROC TREE DATA=TREE NCL=3 OUT=OUT NOPRINT;
ID GROUP;
COPY SLENGTH SWIDTH PLENGTH PWIDTH; } Requests the 2x2 table
PROC FREQ; } showing misclassifications
TABLE CLUSTER*GROUP;
PROC CANDISC NOPRINT OUT=CAN;
CLASS CLUSTER;
VAR SLENGTH SWIDTH PLENGTH PWIDTH; } This series of commands is
PROC PLOT; } used to display cluster
PLOT CAN2*CAN1=CLUSTER; } results. The CANDISC pro-
PROC PLOT; } cedure is run to produce
PLOT CAN2*CAN1=GROUP; } canonical variables for the
 } cluster groups. The first
 } 2 canonical variables are
 } then plotted to show cluster
 } membership

CL-3-Fisher (EML method from SAS)

Same control language as for 2) above except substitute EML for WARD on PROC CLUSTER line.

CL-1-Habbema

Initial partition as specified by input

1	1	2	1	1	1	1	1	1	1
2	1	1	1	1	1	1	2	1	1
1	1	1	1	1	1	1	1	1	1
2	2	1	2	2	1	1	1	1	2
1	2	1	1	1	1	1	2	1	2
2	1	1	1	2	1	1	2	1	2
2	1	2	2	1	1	2	1	1	1
1	2	1	1	2					

Initial group allocation for each observation. The entry for row 1 column 1 refers to observation 1, row 1 column 2 refers to observation 2, and so on

Estimated mean (as a row vector) for each group

ACTIVITY	ANTIGEN
-0.221538	-0.032402 = GROUP 1
-0.282643	-0.040757 = GROUP 2

Group means for each variable based on initial group allocation above

Estimated covariance matrix for group 1 = S_{ij} (group 1)

$$0.031661 = S_1^2$$

$$0.010517 = S_{12} \quad 0.019972 = S_2^2$$

Covariance matrices for each group based on initial group allocation

Estimated covariance matrix for group 2 = S_{ij} (group 2)

$$0.022859 = S_1^2$$

$$0.016834 = S_{12} \quad 0.030533 = S_2^2$$

Proportion from each group as specified by input = Number initially assigned to group i/total number of observations

In loop 55 log likelihood is 77.035

77.035 is the solution of the likelihood equation based on 55 iterations of the EM algorithm

Estimate of mixing proportion for each group
0.508 0.492

Estimate of the final proportion for each group under the normal mixture model

Entity: Final estimates of posterior probabilities of group membership

OBSERVATION	GROUP=1	GROUP=2
1	0.999	0.001
2	0.971	0.029
3	0.245	0.755
4	0.985	0.015
5	0.710	0.290
6	0.986	0.014
7	0.773	0.227
8	0.983	0.017
9	0.958	0.042
10	0.992	0.008
11	0.001	0.999
12	0.983	0.017
13	0.999	0.001
14	0.997	0.003
15	0.948	0.052

These estimates indicate the degree of certainty with which each observation belongs to one of the two groups

For example, observation 1 has a probability of .999 of belonging to group 1 and .001 of belonging to group 2

CL-1-Habbema

16	0.998	0.002
17	0.014	0.986
18	0.966	0.034
19	0.999	0.001
20	0.855	0.145
21	0.957	0.043
22	1.000	0.000
23	0.982	0.018
24	0.976	0.024
25	0.922	0.078
26	0.999	0.001
27	0.899	0.101
28	0.992	0.008
29	0.882	0.118
30	0.993	0.007
31	0.003	0.997
32	0.185	0.815
33	0.001	0.999
34	0.006	0.994
35	0.949	0.051
36	0.000	1.000
37	0.002	0.998
38	0.012	0.988
39	0.223	0.777
40	0.008	0.992
41	0.004	0.996
42	0.045	0.955
43	0.002	0.998
44	0.008	0.992
45	0.274	0.726
46	0.000	1.000

47	0.125	0.875
48	0.000	1.000
49	0.004	0.996
50	0.092	0.908
51	0.606	0.394
52	0.015	0.985
53	0.001	0.999
54	0.620	0.380
55	0.736	0.264
56	0.034	0.966
57	0.591	0.409
58	0.152	0.848
59	0.032	0.968
60	0.899	0.101
61	0.241	0.759
62	0.975	0.025
63	0.970	0.030
64	0.944	0.056
65	0.426	0.574
66	0.636	0.364
67	0.963	0.037
68	0.089	0.911
69	0.992	0.008
70	0.010	0.990
71	0.016	0.984
72	0.126	0.874
73	0.073	0.927
74	0.031	0.969
75	0.000	1.000

Resulting partition of the entities into NG groups

1	1	2	1	1	1	1	1	1	1
2	1	1	1	1	1	2	1	1	1
1	1	1	1	1	1	1	1	1	1
2	2	2	2	1	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2
1	2	2	1	1	2	1	2	2	1
2	1	1	1	2	1	1	2	1	2
2	2	2	2	2					

Final group
allocations after
55 iterations of
clustering algorithm

Number assigned to each group

39 36

Estimates of correct allocation rates for each group

0.934 0.908

Overall estimate of degree of certainty
with which observations are allocated
to each group

Estimate of overall correct allocation rate 0.921 Weighted average
of estimates of correct
allocation rates for each
group

Estimated mean (as a row vector) for each group

ACTIVITY	ANTIGEN
-0.115406	-0.024497 = GROUP 1
-0.365950	-0.045323 = GROUP 2

Group means for each variable
based on final estimates of
posterior probability of group
membership

Estimated covariance matrix for group 1 = S_{ij} (group 1)

0.011245	
0.006548	0.012367

} Based on final
estimates of
posterior prob-
ability of group
membership

Estimated covariance matrix for group 2 = S_{ij} (group 2)

0.015898	
0.015029	0.032278

CL-1-FISHER

Initial partition as specified by input

1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1
2	2	2	2	2	2	2	2	2	2
2	2	3	2	2	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2
3	3	3	3	3	3	3	3	3	3
3	3	2	3	2	3	2	2	2	3
2	3	3	3	2	2	2	3	2	3
3	3	2	3	3	2	3	2	2	3
3	3	3	3	3	3	2	3	2	3

Initial group allocation for
each observation. The entry
for row 1 column 1 refers to
observation 1, row 1 column 2
refers to observation 2, and
so on

Estimated mean (as a row vector) for each group

SLENGTH	SWIDTH	PLENGTH	PWIDTH	
5.005994	3.427995	1.461996	0.246000	GROUP 1
5.920269	2.751557	4.420300	1.434370	GROUP 2
6.869439	3.086106	5.769438	2.105549	GROUP 3

Group means for
each variable
based on initial
group allocation
above

Estimated covariance matrix for group	$1 = S_{ij}(\text{group } 1)$
0.124213	
0.099176	0.143674
0.016347	0.011713
0.010327	0.009296
	0.030165
	0.006070
	0.011106
Estimated covariance matrix for group	$2 = S_{ij}(\text{group } 2)$
0.227175	
0.066786	0.087267
0.141501	0.053037
0.034401	0.028532
	0.277231
	0.117393
	0.085792
Estimated covariance matrix for group	$3 = S_{ij}(\text{group } 3)$
0.241609	
0.016371	0.082387
0.185024	0.011265
-0.008398	0.027246
	0.230741
	0.009312
	0.059419

Covariance
matrices for
each group
based on
initial group
allocation

Estimated common covariance matrix

0.196290			
0.065579	0.104907		
0.110146	0.029316	0.183807	
0.016186	0.021814	0.054552	0.054618

In this run we
specified that KMM
assume equal cov-
ariance matrices
for each group

This is the pooled estimate of that
matrix based on the weighted
average of the individual estimated
covariance matrices

CL-1-FISHER

Proportion from each group as specified by input

0.333 0.427 0.240

.333 = 50/150 = Number initially
assigned to group 1/total number
of observations

In loop 30 log likelihood is

-256.354 = Solution to the likelihood
equation based on 30 iter-
ations of the EM algorithm

Estimate of mixing proportion for each group

0.333 0.330 0.337

Estimate of the final propor-
tion for each group under the
normal mixture model

Entity: Final estimates of posterior probabilities of group membership

OBSERVATION	GROUP 1	GROUP 2	GROUP 3
1	1.000	0.000	0.000
2	1.000	0.000	0.000
3	1.000	0.000	0.000
4	1.000	0.000	0.000
5	1.000	0.000	0.000
6	1.000	0.000	0.000
7	1.000	0.000	0.000
8	1.000	0.000	0.000
9	1.000	0.000	0.000
10	1.000	0.000	0.000
11	1.000	0.000	0.000
12	1.000	0.000	0.000
13	1.000	0.000	0.000
14	1.000	0.000	0.000
15	1.000	0.000	0.000
16	1.000	0.000	0.000
17	1.000	0.000	0.000
18	1.000	0.000	0.000
19	1.000	0.000	0.000
20	1.000	0.000	0.000
21	1.000	0.000	0.000
22	1.000	0.000	0.000
23	1.000	0.000	0.000
24	1.000	0.000	0.000
25	1.000	0.000	0.000
26	1.000	0.000	0.000
27	1.000	0.000	0.000
28	1.000	0.000	0.000
29	1.000	0.000	0.000
30	1.000	0.000	0.000
31	1.000	0.000	0.000
32	1.000	0.000	0.000
33	1.000	0.000	0.000
34	1.000	0.000	0.000
35	1.000	0.000	0.000
36	1.000	0.000	0.000
37	1.000	0.000	0.000
38	1.000	0.000	0.000
39	1.000	0.000	0.000
40	1.000	0.000	0.000
41	1.000	0.000	0.000
42	1.000	0.000	0.000
43	1.000	0.000	0.000
44	1.000	0.000	0.000
45	1.000	0.000	0.000
46	1.000	0.000	0.000
47	1.000	0.000	0.000
48	1.000	0.000	0.000

These estimates indicate the degree of certainty with which each observation belongs to one of the three groups. Observation 1 has a probability of 1.0 of belonging to group 1 and 0 of belonging to the other two groups

49	1.000	0.000	0.000
50	1.000	0.000	0.000
51	0.000	0.999	0.001
52	0.000	1.000	0.000
53	0.000	1.000	0.000
54	0.000	1.000	0.000
55	0.000	1.000	0.000
56	0.000	1.000	0.000
57	0.000	0.999	0.001
58	0.000	1.000	0.000
59	0.000	1.000	0.000
60	0.000	1.000	0.000
61	0.000	1.000	0.000
62	0.000	1.000	0.000
63	0.000	0.704	0.296
64	0.000	1.000	0.000
65	0.000	1.000	0.000
66	0.000	1.000	0.000
67	0.000	1.000	0.000
68	0.000	1.000	0.000
69	0.000	1.000	0.000
70	0.000	0.997	0.003
71	0.000	1.000	0.000
72	0.000	0.967	0.033
73	0.000	1.000	0.000
74	0.000	1.000	0.000
75	0.000	1.000	0.000
76	0.000	0.998	0.002
77	0.000	0.999	0.001
78	0.000	0.127	0.873
79	0.000	1.000	0.000
80	0.000	0.999	0.001
81	0.000	0.979	0.021
82	0.000	0.133	0.867
83	0.000	0.868	0.132
84	0.000	0.991	0.009
85	0.000	1.000	0.000
86	0.000	1.000	0.000
87	0.000	0.988	0.012
88	0.000	0.997	0.003
89	0.000	0.998	0.002
90	0.000	0.994	0.006
91	0.000	1.000	0.000
92	0.000	0.999	0.001
93	0.000	0.929	0.071
94	0.000	0.979	0.021
95	0.000	0.999	0.001
96	0.000	1.000	0.000
97	0.000	1.000	0.000
98	0.000	1.000	0.000
99	0.000	1.000	0.000

100	0.000	1.000	0.000
101	0.000	0.000	1.000
102	0.000	0.000	1.000
103	0.000	0.000	1.000
104	0.000	0.000	1.000
105	0.000	0.148	0.852
106	0.000	0.000	1.000
107	0.000	0.000	1.000
108	0.000	0.000	1.000
109	0.000	0.000	1.000
110	0.000	0.000	1.000
111	0.000	0.000	1.000
112	0.000	0.002	0.998
113	0.000	0.000	1.000
114	0.000	0.000	1.000
115	0.000	0.009	0.991
116	0.000	0.000	1.000
117	0.000	0.001	0.999
118	0.000	0.094	0.906
119	0.000	0.123	0.877
120	0.000	0.003	0.997
121	0.000	0.162	0.838
122	0.000	0.001	0.999
123	0.000	0.000	1.000
124	0.000	0.004	0.996
125	0.000	0.001	0.999
126	0.000	0.089	0.911
127	0.000	0.302	0.698
128	0.000	0.000	1.000
129	0.000	0.000	1.000
130	0.000	0.000	1.000
131	0.000	0.000	1.000
132	0.000	0.000	1.000
133	0.000	0.746	0.254
134	0.000	0.002	0.998
135	0.000	0.000	1.000
136	0.000	0.073	0.927
137	0.000	0.000	1.000
138	0.000	0.006	0.994
139	0.000	0.022	0.978
140	0.000	0.000	1.000
141	0.000	0.000	1.000
142	0.000	0.000	1.000
143	0.000	0.001	0.999
144	0.000	0.005	0.995
145	0.000	0.000	1.000
146	0.000	0.000	1.000
147	0.000	0.000	1.000
148	0.000	0.000	1.000
149	0.000	0.000	1.000
150	0.000	0.008	0.992

CL-1-FISHER

Resulting partition of the entities into NG groups

1	1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1	1
1	1	1	1	1	1	1	1	1	1	1
2	2	2	2	2	2	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2	2
2	2	2	2	2	2	2	3	2	2	2
2	3	2	2	2	2	2	2	2	2	2
2	2	2	2	2	2	2	2	2	2	2
3	3	3	3	3	3	3	3	3	3	3
3	3	3	3	3	3	3	3	3	3	3
3	3	3	3	3	3	3	3	3	3	3
3	3	2	3	3	3	3	3	3	3	3
3	3	3	3	3	3	3	3	3	3	3

Final group allocations after
30 iterations

Number assigned to each group

50 49 51

Estimates of correct allocation rates for each group

1.000 0.973 0.983

Overall estimate
of degree of certainty
with which observations
are allocated to each
group

Estimate of overall correct allocation rate 0.985 = Weighted average of
estimates of correct
allocation rates for
each group

Estimated mean (as a row vector) for each group

SLENGTH	SWIDTH	PLENGTH	PWIDTH
5.005994	3.427995	1.461996	0.246000
5.942309	2.760773	4.258801	1.319220
6.574652	2.980818	5.539058	2.024963

Group means for
each variable based
on estimates of
posterior proba-
bility of group membership

Estimated common covariance matrix

0.263932		
0.089847	0.111946	
0.169658	0.051118	0.186544
0.039336	0.029976	0.041973

This pooled estimate of the
common covariance matrix is
based on the final estimates
of posterior probability of
group membership
0.039709

SAS

CL-2-Habbema

WARD'S MINIMUM VARIANCE CLUSTER ANALYSIS

EIGENVALUES OF THE COVARIANCE MATRIX

EIGENVALUE	DIFFERENCE $(\lambda_i - \lambda_{i+1})$	PROPORTION $\lambda_i / \sum \lambda_i$	CUMULATIVE
1 0.038779	0.025371	0.743072	0.74307
2 0.013409		0.256928	1.00000

ROOT-MEAN-SQUARE TOTAL-SAMPLE STANDARD DEVIATION = 0.161536

ROOT-MEAN-SQUARE DISTANCE BETWEEN OBSERVATIONS = 0.323072 → squared Euclidean distances

NUMBER = number of clusters
OF
formed at each step
CLUSTERS CLUSTERS JOINED

FREQUENCY = number of observations in the newly formed cluster
OF NEW
CLUSTER SEMIPARTIAL
R-SQUARED

Clusters joined identifies the two clusters being joined at a particular step. Clusters of one observation being joined is identified as OBn, where n is the observation number. Clusters of more than one observation are identified as CLn, where n is the number of clusters existing after the cluster is formed.

R-squared is the decrease in the proportion of variance accounted for resulting from joining two clusters.

.000007 is the decrease in the proportion of variances accounted for resulting from joining observations 4 and 6 into one cluster.

R-squared is the squared multiple correlation and
is the proportion of variances accounted for by the clusters at a particular step.

i.e. .997537 = R² for 49 clusters

CL70	OB46	0.998967	0.000122
57	OB14	0.998835	0.000132
56	OB17	0.998691	0.000143
55	OB31	0.998537	0.000154
54	CL71	0.998371	0.000166
53	OB23	0.998186	0.000185
52	OB40	0.997997	0.000189
51	OB2	0.997808	0.000190
50	OB5	0.997537	0.000271
49	OB61	0.997256	0.000281
48	OB75	0.997056	0.000290
47	CL74	0.996956	0.000305
46	OB41	0.996661	0.000323
45	OB58	0.996338	0.000334
44	OB1	0.996005	0.000353
43	OB56	0.995652	0.000375
42	CL59	0.995277	0.000426
41	CL62	0.994851	0.000429
40	CL72	0.994421	0.000444
39	CL56	0.993977	0.000474
38	CL50	0.993504	0.000491
37	CL67	0.993023	0.000493
36	OB65	0.992530	0.000532
35	CL47	0.991998	0.000564
34	OB10	0.991434	0.000590
33	CL69	0.990844	0.000628
32	CL53	0.990215	0.000646
31	CL55	0.989659	0.000773
30	OB70	0.988797	0.001050
29	CL54	0.987747	0.001088
28	CL40	0.986658	0.001117
27	CL58	0.985541	0.001243
26	CL44	0.984298	0.001263
25	OB60	0.983035	0.001421
24	CL48	0.981613	0.001472
23	CL45	0.979741	0.001980
22	CL38	0.977762	0.002592
21	CL41	0.975170	0.002592

	CL51	0.002801	0.972369
20	CL32	7	5
	CL42	5	0.002858
19	CL36	5	0.969511
18	CL39	6	0.003321
17	CL23	6	0.966190
16	CL27	12	0.003847
15	0B11	12	0.962343
14	CL21	8	0.005464
13	CL15	8	0.956879
12	CL17	2	0.005975
11	CL18	2	0.950904
10	CL24	9	0.007694
9	CL26	9	0.943210
8	CL16	5	0.008771
7	CL9	15	0.934439
6	CL20	15	0.009699
5	CL14	8	0.924741
4	CL6	8	0.009889
3	CL7	8	0.914851
2	CL3	9	0.010854
1	CL2	9	0.903998
		9	0.013422
		13	0.890575
		13	0.018916
		24	0.871660
		24	0.025938
		16	0.845722
		16	0.029542
		22	0.816180
		22	0.040895
		21	0.775285
		21	0.057587
		32	0.717698
		32	0.095677
		54	0.622021
		54	0.136697
		75	0.485324
			0.000000

SAS

CL-2-Habbema

TABLE OF CLUSTER BY GROUP

CLUSTER	GROUP							TOTAL
		FREQUENCY						
		PERCENT						
		ROW PCT						
		COL PCT						
			1;		2;			
Cell frequency = 27			1	27	27	54	= row total	
Cell percent = .36			1	36.00	36.00	72.00	= row percent	
Row percent = .50				50.00	50.00		= 54/75	
column percent = .90				90.00	60.00		= .72	
			2	3	18	21		
				4.00	24.00	28.00		
				14.29	85.71			
				10.00	40.00			
			TOTAL	30	45	75		
				40.00	60.00	100.00		
							column total = 30	
							column percent = 30/75 = .4	

This 2x2 table shows misclassifications. GROUP contains the true group allocation of each observation and CLUSTER contains the group allocation based on the Ward's clustering. 27 observations were correctly assigned to cluster 1 and 18 to cluster 2. 27 observations from group 2 were incorrectly assigned to cluster 1 while 3 from group 1 were incorrectly assigned to cluster 2.

WARD'S MINIMUM VARIANCE CLUSTER ANALYSIS
EIGENVALUES OF THE COVARIANCE MATRIX

EIGENVALUE	DIFFERENCE $(\lambda_1 - \lambda_{i+1})$	PROPORTION $\lambda_1 / \sum \lambda_i$	CUMULATIVE
1 4.22824	3.98657	0.924619	0.92462
2 0.24267	0.16446	0.053066	0.97769
3 0.07821	0.05437	0.017103	0.99479
4 0.02384	.	0.005212	1.00000

ROOT-MEAN-SQUARE TOTAL-SAMPLE STANDARD DEVIATION = 1.06922
ROOT-MEAN-SQUARE DISTANCE BETWEEN OBSERVATIONS = 3.02422

NUMBER OF CLUSTERS	CLUSTERS JOINED	FREQUENCY OF NEW CLUSTER	SEMIPARTIAL R-SQUARED	R-SQUARED
149 OB117	OB125	2	0.000000	1.000000
148 OB14	OB46	2	0.000007	0.999993
147 OB6	OB48	2	0.000007	0.999985
146 OB1	OB50	2	0.000007	0.999978
145 OB137	OB149	2	0.000007	0.999971
144 OB17	OB40	2	0.000007	0.999963
143 OB52	OB62	2	0.000015	0.999949
142 OB5	OB38	2	0.000015	0.999934
141 OB124	OB144	2	0.000015	0.999919
140 OB119	OB126	2	0.000015	0.999905
139 OB65	OB97	2	0.000015	0.999890
138 OB53	OB54	2	0.000015	0.999875
137 OB22	OB29	2	0.000015	0.999861
136 OB23	OB35	2	0.000015	0.999846
135 OB11	OB16	2	0.000015	0.999831
134 OB76	OB90	2	0.000015	0.999817
133 OB59	OB69	2	0.000015	0.999802

.000007 is the decrease in the proportion
of variance explained resulting from
joining observations 137 and 149. The
associated R² is .999971

0.999787	0.999773	0.999758	0.999743	0.999721	0.999700	0.999679	0.999669	0.999655	0.999633	0.999609	0.999584	0.999555	0.999525	0.999496	0.999467	0.999435	0.999401	0.999366	0.999330	0.999293	0.999249	0.999205	0.999161	0.999117	0.999073	0.999029	0.999090	0.998929	0.998877	0.998826	0.998775	0.998723	0.998669	0.998616	0.998560	0.998501								
OB61	OB49	OB47	OB44	OB121	OB12	OB122	OB85	CL149	CL135	CL138	CL111	OB64	OB9	OB118	CL146	OB112	OB86	OB96	OB72	OB93	OB13	OB66	CL147	CL142	CL137	CL134	OB114	OB110	CL143	OB123	OB88	CL119	OB84	OB51	OB106	OB75	OB56	OB107	CL130	CL141	CL127	OB42	CL132	OB82
2	2	2	2	2	2	3	2	3	3	3	2	2	3	2	2	2	2	2	2	2	2	3	3	3	3	3	2	2	2	2	2	2	2	2	2	3	3	3						
132	131	130	129	128	127	126	125	124	123	122	121	120	119	118	117	116	115	114	113	112	111	110	109	108	107	106	105	104	103	102	101	100	99	98	97	96	95							

94	OB2	CL129	OB20	2	2	0.000064	0.998310
93	OB10	CL149	OB21	3	3	0.000066	0.998244
92	OB113	CL124	OB22	8	8	0.000068	0.998176
91	CL98	OB92	OB23	2	2	0.000073	0.998103
90	OB89	CL117	OB24	6	6	0.000073	0.998030
89	CL102	OB80	OB25	3	3	0.000076	0.997956
88	OB78	OB133	OB26	2	2	0.000081	0.997880
87	OB102	CL113	OB27	3	3	0.000081	0.997800
86	OB26	OB36	OB28	2	2	0.000081	0.997719
85	OB26	CL94	OB18	4	4	0.000083	0.997638
84	OB26	CL122	CL125	5	5	0.000086	0.997555
83	OB26	OB105	OB120	2	2	0.000088	0.997470
82	OB28	OB28	OB45	2	2	0.000088	0.997382
81	OB28	CL145	OB145	3	3	0.000091	0.997293
80	OB28	OB58	CL112	4	4	0.000092	0.997111
79	OB28	CL95	OB115	4	4	0.000094	0.997017
78	CL95	CL128	OB138	3	3	0.000095	0.996922
77	CL128	CL92	OB148	4	4	0.000097	0.996825
76	CL92	CL144	OB119	3	3	0.000100	0.996725
75	CL144	OB57	CL107	3	3	0.000103	0.996622
74	OB57	CL110	OB128	3	3	0.000103	0.996519
73	CL110	OB77	OB99	2	2	0.000117	0.996402
72	OB77	CL91	CL136	10	10	0.000121	0.996281
71	CL91	CL87	OB83	3	3	0.000125	0.996156
70	CL87	OB104	OB135	2	2	0.000125	0.996031
69	OB104	CL86	CL108	5	5	0.000126	0.995905
68	CL86	CL100	CL115	6	6	0.000139	0.995766
67	CL100	CL133	OB67	3	3	0.000142	0.995624
66	CL133	CL75	OB34	4	4	0.000142	0.995482
65	CL75	OB89	OB7	7	7	0.000142	0.995340
64	OB89	OB55	OB98	2	2	0.000147	0.995193
63	OB55	CL105	OB87	3	3	0.000154	0.995039
62	CL105	OB63	CL114	3	3	0.000159	0.994880
61	OB63	CL116	OB15	4	4	0.000160	0.994720
60	CL116	OB112	CL99	4	4	0.000170	0.994550
59	OB112	CL109	CL118	5	5	0.000170	0.994380
58	CL109	CL82	OB41	3	3	0.000176	0.994204
57	CL82						

56	CL126	CL111	0.000194
55	CL72	CL121	4
54	CL74	CL139	4
53	CL101	OB132	5
52	OB25	OB39	5
51	CL120	CL90	3
50	CL70	CL77	2
49	CL65	CL97	2
48	CL76	OB129	4
47	CL66	OB71	4
46	CL106	CL93	5
45	CL52	CL81	4
44	CL58	OB95	6
43	CL57	CL104	5
42	CL96	OB100	4
41	CL49	CL85	8
40	CL64	CL67	13
39	CL103	OB127	3
38	OB101	CL73	4
37	CL90	OB147	4
36	CL83	CL51	9
35	CL88	CL54	8
34	CL59	CL37	8
33	OB103	OB110	2
32	CL61	CL56	7
31	CL44	CL63	8
30	CL84	OB37	5
29	CL60	CL41	12
28	CL79	CL55	8
27	CL71	CL46	15
26	CL31	CL42	12
25	CL50	OB136	7
24	CL62	CL78	7
23	CL38	CL68	9
22	CL30	OB30	6
21	CL33	CL69	4
20	CL36	OB139	10
19	CL22	CL40	19

18	CL39	CL25	0.001249
17	CL29	CL45	0.975524
16	CL32	CL34	0.001351
15	CL24	CL28	0.974172
14	CL21	CL53	0.001462
13	CL18	CL48	0.972710
12	CL16	CL23	0.001641
11	CL14	CL43	0.971069
10	CL26	CL20	0.969196
9	CL27	CL17	0.001873
8	CL15	CL35	0.966925
7	CL10	CL47	0.002271
6	CL8	CL13	0.964661
5	CL9	CL19	0.002500
4	CL12	CL11	0.962151
3	CL6	CL7	0.002694
2	CL3	CL4	0.959457
1	CL5	CL2	0.003060
10			0.956397
			0.953302
			0.003095
			0.005811
			0.947491
			0.006042
			0.941449
			0.010532
			0.930917
			0.913673
			0.017245
			0.883621
			0.030051
			0.1111026
			0.772595
			0.000000
			0.772595

SAS

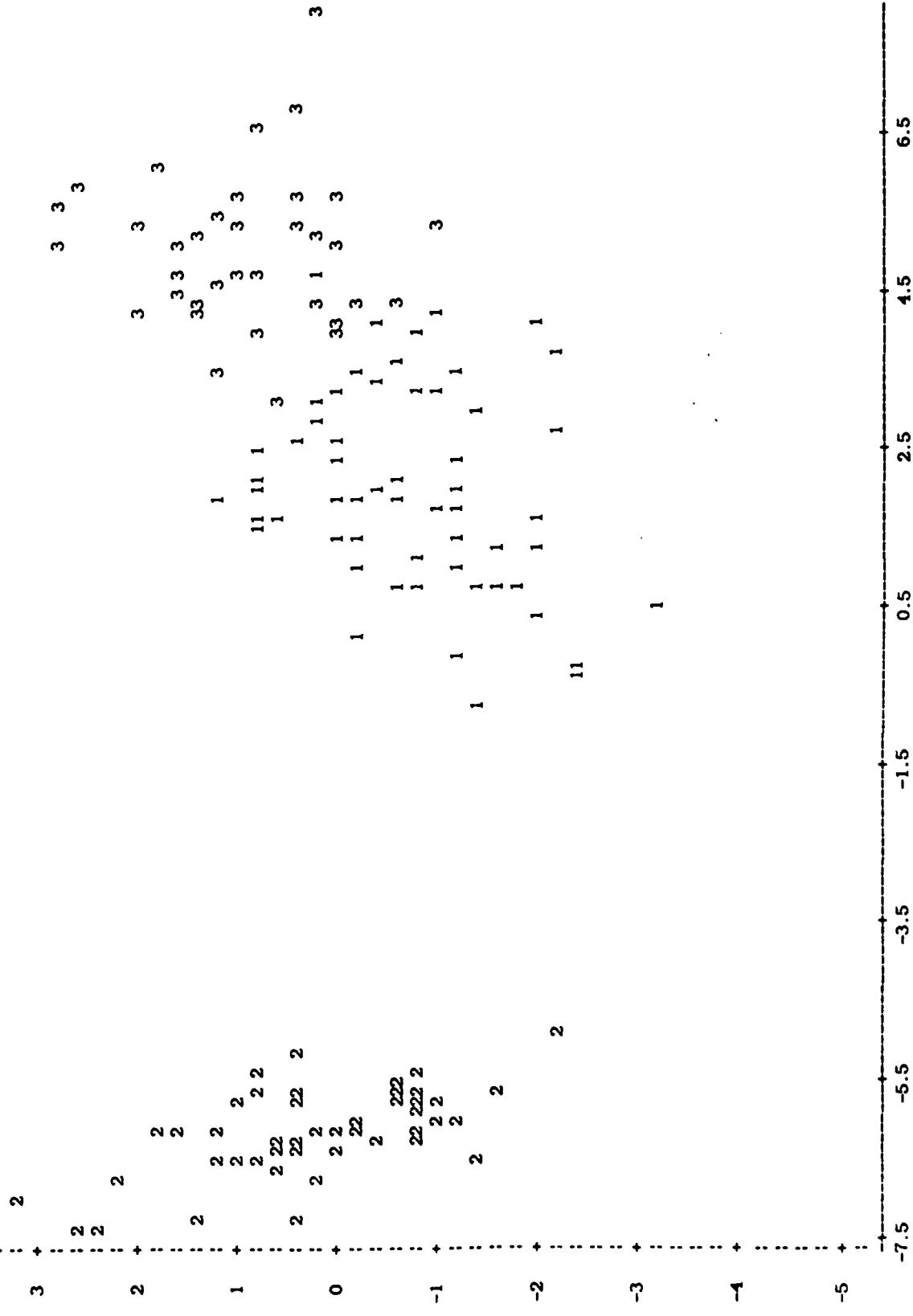
TABLE OF CLUSTER BY GROUP

CLUSTER	GROUP	FREQUENCY			TOTAL
		1	2	3	
1	0.00	49	15	64	64
	0.00	32.67	10.00	42.67	
	0.00	76.56	23.44	0.00	
2	0.00	98.00	30.00	0.00	50
	50	0	0	0.00	
	33.33	0.00	0.00	33.33	
3	100.00	0.00	0.00	0.00	36
	100.00	0.00	0.00	0.00	
	0.00	0.00	0.00	0.00	
TOTAL		50	50	50	150
		33.33	33.33	33.33	100.00

2x2 table displaying misclassifications. The true group allocation is labelled GROUP and is displayed as columns. The rows show the group allocation based on Ward's Clustering. One observation from GROUP 2 was misclassified and 15 from GROUP 3

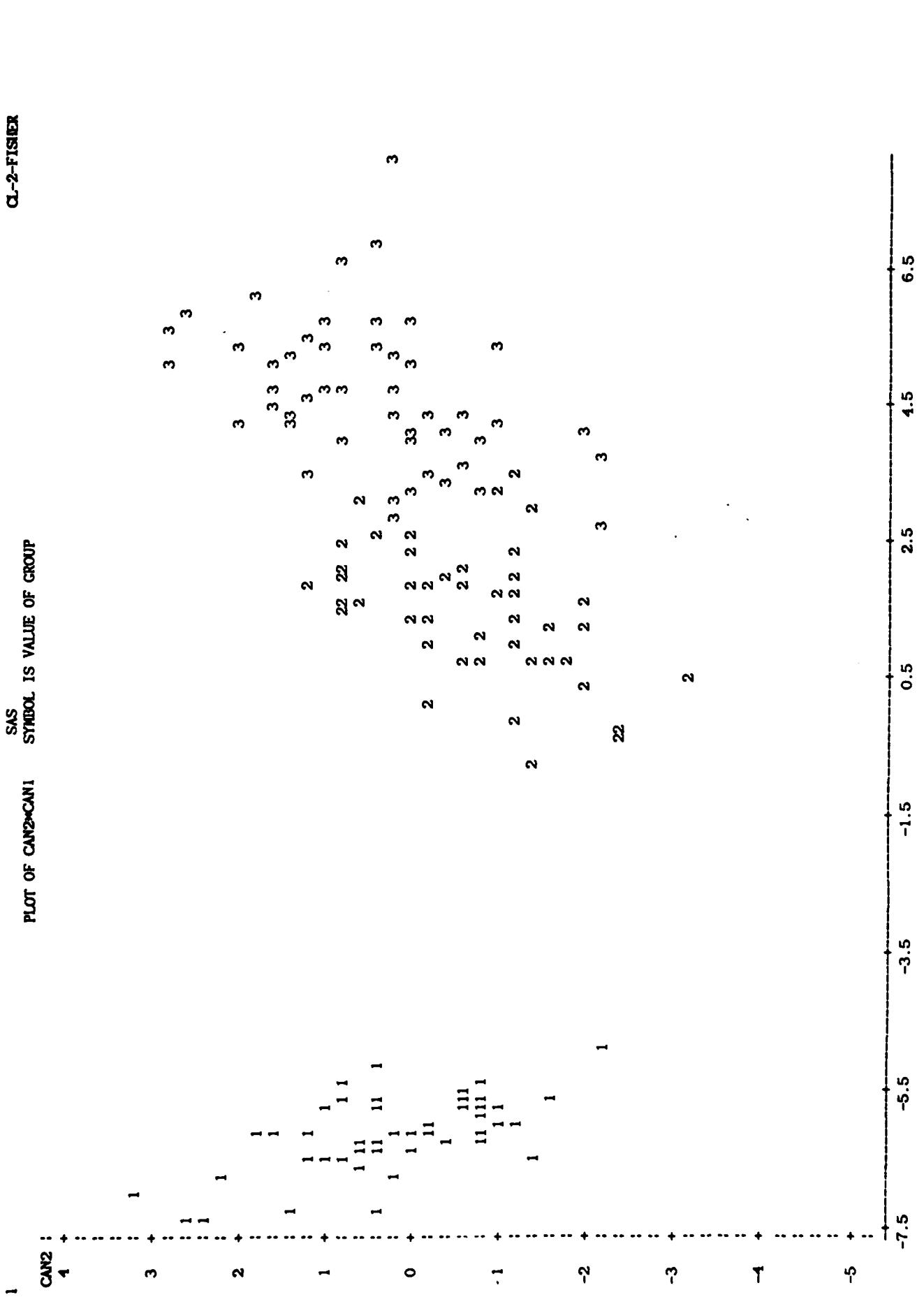
CL-2-FISHER

PLOT OF CAN2*CAN1
SAS SYMBOL IS VALUE OF CLUSTER



NOTE: 13 OBS HIDDEN

This plot displays the results when three clusters are formed. The first two canonical variables (CAN1 and CAN2) for discriminating among the three clusters were computed and plotted to show cluster membership. The symbol plotted is the value of CLUSTER.



NOTE: 13 OBS HIDDEN
This plot is exactly the same as the one on the previous page except the symbol plotted is the value of GROUP.

SAS
 EQUAL VARIANCE MAXIMUM LIKELIHOOD METHOD
 EIGENVALUES OF THE COVARIANCE MATRIX

	EIGENVALUE	DIFFERENCE	PROPORTION	CUMULATIVE
1	0.038779	0.025371	0.743072	0.74307
2	0.013409	.	0.25628	1.00000

ROOT-MEAN-SQUARE TOTAL-SAMPLE STANDARD DEVIATION = 0.161536
 ROOT-MEAN-SQUARE DISTANCE BETWEEN OBSERVATIONS = 0.323072

NUMBER OF CLUSTERS	CLUSTERS JOINED	FREQUENCY OF NEW CLUSTER	LOG LIKELIHOOD RATIO	LOG LIKELIHOOD
74	OB4	OB6	2	1692.3
73	OB20	OB29	2	1498.5
72	OB49	OB52	2	121.7
71	OB3	OB34	2	1376.9
70	OB8	OB12	2	1311.4
69	OB9	OB21	2	1264.4
68	OB15	OB25	2	1227.1
67	OB36	OB39	2	37.3953
66	OB35	OB50	2	46.9378
65	OB43	OB44	2	65.4659
64	OB28	OB30	2	193.8
63	OB7	OB37	2	28.3152
62	OB18	OB45	2	24.8173
61	OB54	OB71	2	1141.7
60	OB64	OB67	2	24.1433
59	OB63	OB73	2	1117.5
58	CL63	OB38	2	20.5569
57	CL70	OB46	2	1097.0
56	OB14	OB47	2	20.0080
55	CL71	CL66	4	1077.0
				1042.8
				1028.1
				999.5
				969.3
				954.9

942.3	12.5620
930.2	12.1489
917.8	12.3538
905.7	12.0734
894.9	10.7887
883.1	11.8875
870.3	12.7666
858.5	11.8045
847.9	10.5419
837.9	10.0196
827.6	10.3168
818.0	9.6231
809.6	8.4188
802.3	7.3100
795.2	7.0361
788.5	6.6825
781.7	6.7983
775.3	6.4054
769.2	6.1010
762.9	6.3651
756.7	6.1221
751.1	5.6129
745.6	5.5075
739.8	5.7742
732.9	6.9427
726.0	7.6944
718.3	7.4183
710.8	5.8181
705.0	5.6233
699.4	6.3334
693.1	5.9405
687.2	10.9017
676.4	6.1111
668.3	7.5983
660.7	6.8347
653.9	6.1380
647.7	5.3666
OB32	2
OB55	2
CL64	3
OB48	2
OB24	2
CL57	5
OB69	2
OB75	2
CL65	3
OB41	3
CL60	3
OB58	2
OB19	2
CL59	3
OB68	3
CL73	4
CL62	4
CL72	4
CL61	4
OB56	2
CL58	4
OB26	3
OB32	6
CL50	3
CL56	3
CL49	3
OB33	6
OB59	3
CL52	4
OB72	2
OB42	5
CL55	3
CL55	3
CL69	3
OB27	3
CL54	3
CL36	6
OB70	2
OB57	5
CL41	5
CL53	5
CL27	6
CL39	8
CL42	8
OB13	3
CL44	3
CL44	3
OB60	3
CL47	3
CL36	12
CL25	22
CL21	21
CL28	20
OB22	19
CL34	2
CL38	6
CL46	17

16	CL37	CL48	8	4.1815	638.2
15	CL30	CL40	8	9.5200	628.7
14	CL22	CL32	15	13.1016	615.6
13	CL16	CL17	14	9.5519	606.0
12	CL21	CL26	13	7.5668	598.5
11	0811	0851	2	7.8476	590.6
10	CL11	CL31	5	7.6117	583.0
9	CL14	CL15	23	5.4097	577.6
8	CL20	CL18	12	9.1725	568.4
7	CL24	CL9	26	5.7861	562.6
6	CL13	CL19	16	6.6930	555.9
5	CL12	CL23	16	4.7377	551.2
4	CL7	CL6	42	6.9678	544.2
3	CL4	CL8	54	-10.1224	554.4
2	CL5	CL10	21	-5.2539	559.6
1	CL3	CL2	75	10.6897	548.9

SAS
TABLE OF CLUSTER BY GROUP

CLUSTER	GROUP	FREQUENCY		PERCENT		TOTAL
		ROW PCT	COL PCT	ROW PCT	COL PCT	
1	1	27	27	54		
	2	36.00	36.00	72.00		
	3	50.00	50.00			
	4	90.00	60.00			
2	1	3	18	21		
	2	4.00	24.00	28.00		
	3	14.29	85.71			
	4	10.00	40.00			
TOTAL		30	45	75		
		40.00	60.00	100.00		

SAS
 EQUAL VARIANCE MAXIMUM LIKELIHOOD METHOD
 EIGENVALUES OF THE COVARIANCE MATRIX

EIGENVALUE	DIFFERENCE	PROPORTION	CUMULATIVE
1 4.22824	3.98557	0.924619	0.92462
2 0.24267	0.16446	0.053066	0.97769
3 0.07821	0.05437	0.017103	0.99479
4 0.02384		0.005212	1.00000

ROOT-MEAN-SQUARE TOTAL-SAMPLE STANDARD DEVIATION = 1.06922
 ROOT-MEAN-SQUARE DISTANCE BETWEEN OBSERVATIONS = 3.02422

NUMBER OF CLUSTERS	CLUSTERS JOINED	FREQUENCY OF NEW CLUSTER	LOG LIKELIHOOD RATIO	LOG LIKELIHOOD
149	OB117	OB125	2	
148	OB14	OB46	2	4016.3
147	OB6	OB48	2	3603.2
146	OB1	OB50	2	3362.7
145	OB137	OB149	2	3192.9
144	OB17	OB40	2	3061.7
143	OB52	OB62	2	2862.6
142	OB5	OB38	2	2714.6
141	OB124	OB144	2	2597.0
140	OB119	OB126	2	2499.5
139	OB65	OB97	2	2416.4
138	OB53	OB54	2	2344.1
137	OB23	OB35	2	2280.1
136	OB22	OB29	2	2222.9
135	OB11	OB16	2	2171.1
134	OB76	OB90	2	2123.8
133	OB59	OB69	2	2080.4
132	OB60	OB61	2	2040.3

OB21	OB49	37.2422
130	OB8	34.7396
129	OB3	32.5317
128	OB9	45.5238
127	CL146	1848.5
126	OB111	1935.8
125	OB118	1890.3
124	OB64	1848.3
123	CL135	OB44
122	CL138	CL148
121	OB86	OB112
120	OB72	OB122
119	OB66	OB121
118	OB13	OB124
117	CL147	OB85
116	CL142	OB85
115	CL136	OB85
114	OB134	OB85
113	OB130	OB85
112	CL134	OB85
111	CL143	OB85
110	OB123	OB85
109	OB114	OB85
108	OB109	OB85
107	OB68	OB85
106	OB4	OB85
105	CL127	OB85
104	OB106	OB85
103	OB107	OB85
102	OB51	OB85
101	OB75	OB85
100	OB56	OB85
99	CL141	OB70
98	CL130	OB143
97	CL105	OB33
96	OB31	OB33
95	CL117	OB106
94	OB82	CL123

93	92	OB2	0874	CL129	3	3	17.3356	908.6
91	OB113	OB113	CL149	CL149	3	3	18.1585	891.3
90	OB10	OB20	OB20	OB20	2	2	17.6839	873.2
89	CL122	CL124	CL124	CL124	5	5	18.7901	855.5
88	CL100	OB80	OB80	OB80	3	3	17.9123	836.7
87	OB89	OB92	OB92	OB92	2	2	17.5339	818.8
86	CL92	OB18	OB18	OB18	4	4	17.7140	801.2
85	OB102	CL113	CL113	CL113	3	3	16.9813	783.5
84	OB78	OB133	OB133	OB133	2	2	17.3308	766.5
83	OB26	OB36	OB36	OB36	2	2	16.6790	749.2
82	OB58	CL112	CL112	CL112	4	4	16.8950	732.5
81	CL145	OB145	OB145	OB145	3	3	16.5465	715.7
80	CL94	OB115	OB115	OB115	4	4	15.9775	699.1
79	CL97	CL137	CL137	CL137	10	10	15.3316	683.1
78	CL91	OB148	OB148	OB148	4	4	14.9811	667.8
77	OB105	OB120	OB120	OB120	2	2	14.4466	652.8
76	OB28	OB45	OB45	OB45	2	2	13.9662	638.4
75	CL125	OB138	OB138	OB138	3	3	13.8027	624.4
74	CL144	OB19	OB19	OB19	3	3	14.1647	610.6
73	CL108	OB128	OB128	OB128	3	3	14.0609	596.5
72	OB857	CL107	CL107	CL107	3	3	14.0609	582.4
71	CL85	CL110	CL110	CL110	5	5	13.5435	568.8
70	CL95	CL115	CL115	CL115	9	9	13.9353	554.9
69	OB777	OB899	OB899	OB899	2	2	15.0862	541.4
68	CL84	OB833	OB833	OB833	3	3	14.5910	526.4
67	OB104	OB135	OB135	OB135	2	2	15.0893	511.8
66	CL74	OB34	OB34	OB34	4	4	15.1889	496.7
65	CL133	OB67	OB67	OB67	3	3	15.2430	481.5
64	CL79	OB7	OB7	OB7	11	11	14.9585	466.2
63	CL111	CL119	CL119	CL119	5	5	14.6028	451.3
62	CL102	OB87	OB87	OB87	3	3	14.8579	436.7
61	CL116	OB15	OB15	OB15	4	4	14.3220	421.8
60	OB55	OB98	OB98	OB98	2	2	13.9645	407.5
59	OB63	CL114	CL114	CL114	3	3	13.8010	393.5
58	OB112	CL99	CL99	CL99	4	4	13.7858	379.7
57	CL77	OB141	OB141	OB141	3	3	14.5532	365.9
56	CL126	CL109	CL109	CL109	4	4	14.1081	351.4
								337.3

55	CL72	CL139	5	13.6412	323.6
54	CL69	CL121	4	13.8841	309.8
53	CL68	CL75	6	14.2400	295.5
52	CL89	CL87	7	13.9381	281.6
51	CL70	CL98	12	13.1141	268.5
50	CL66	CL96	6	13.4041	255.1
49	CL103	OB132	3	12.9165	242.1
48	OB25	OB39	2	14.0625	228.1
47	CL106	CL90	5	13.9022	214.2
46	CL50	CL83	8	14.0355	200.1
45	CL78	OB129	5	13.5881	186.6
44	CL57	CL104	5	13.3811	173.2
43	CL48	CL76	4	12.9790	160.2
42	CL52	CL93	10	12.5952	147.6
41	CL65	OB71	4	12.1714	135.4
40	CL63	OB95	6	11.7183	123.7
39	CL88	CL55	8	11.9019	111.8
38	CL58	CL81	7	12.1729	99.6391
37	CL64	CL47	16	12.5030	87.1362
36	CL61	CL46	12	11.5860	75.5502
35	CL53	OB127	7	12.7557	62.7945
34	OB101	CL71	6	13.2697	49.5248
33	CL59	CL56	7	12.6890	36.8369
32	CL38	OB147	8	12.3602	24.4767
31	CL82	CL54	8	11.9039	12.5727
30	CL40	CL60	8	11.2545	1.3182
29	CL101	OB100	3	12.5860	-11.2678
28	CL42	CL120	12	13.8733	-25.1411
27	CL44	OB110	6	13.8089	-38.9500
26	CL86	CL51	16	13.7295	-52.6795
25	CL26	OB37	17	12.6505	-65.3300
24	CL62	CL31	11	11.9355	-77.2655
23	CL35	OB136	8	12.8183	-90.0638
22	CL34	CL73	9	12.2185	-102.3
21	CL37	CL36	28	10.8425	-113.1
20	CL80	CL45	9	9.8004	-122.9
19	OB103	CL67	3	15.3968	-138.3
18	CL33	CL32	15	15.2276	-153.6

17	CL25	0830	18	15.8760
16	CL23	CL20	17	15.6063
15	CL30	CL28	20	15.1236
14	CL18	CL22	24	12.1927
13	CL15	08139	21	15.9966
12	CL24	CL39	19	17.4915
11	CL13	CL29	24	20.6397
10	CL27	CL49	9	19.4921
9	CL21	CL43	32	17.9355
8	CL19	CL10	12	20.3710
7	CL12	CL16	36	43.3135
6	CL9	CL17	50	39.8843
5	CL11	CL41	28	34.1956
4	CL7	CL14	60	78.4355
3	CL4	CL8	72	163.6
2	CL3	CL5	100	198.1
1	CL6	CL2	150	697.7

-245.9

-266.5

-286.0

-303.9

-324.3

-367.6

-407.5

-441.7

-520.1

-683.7

-881.8

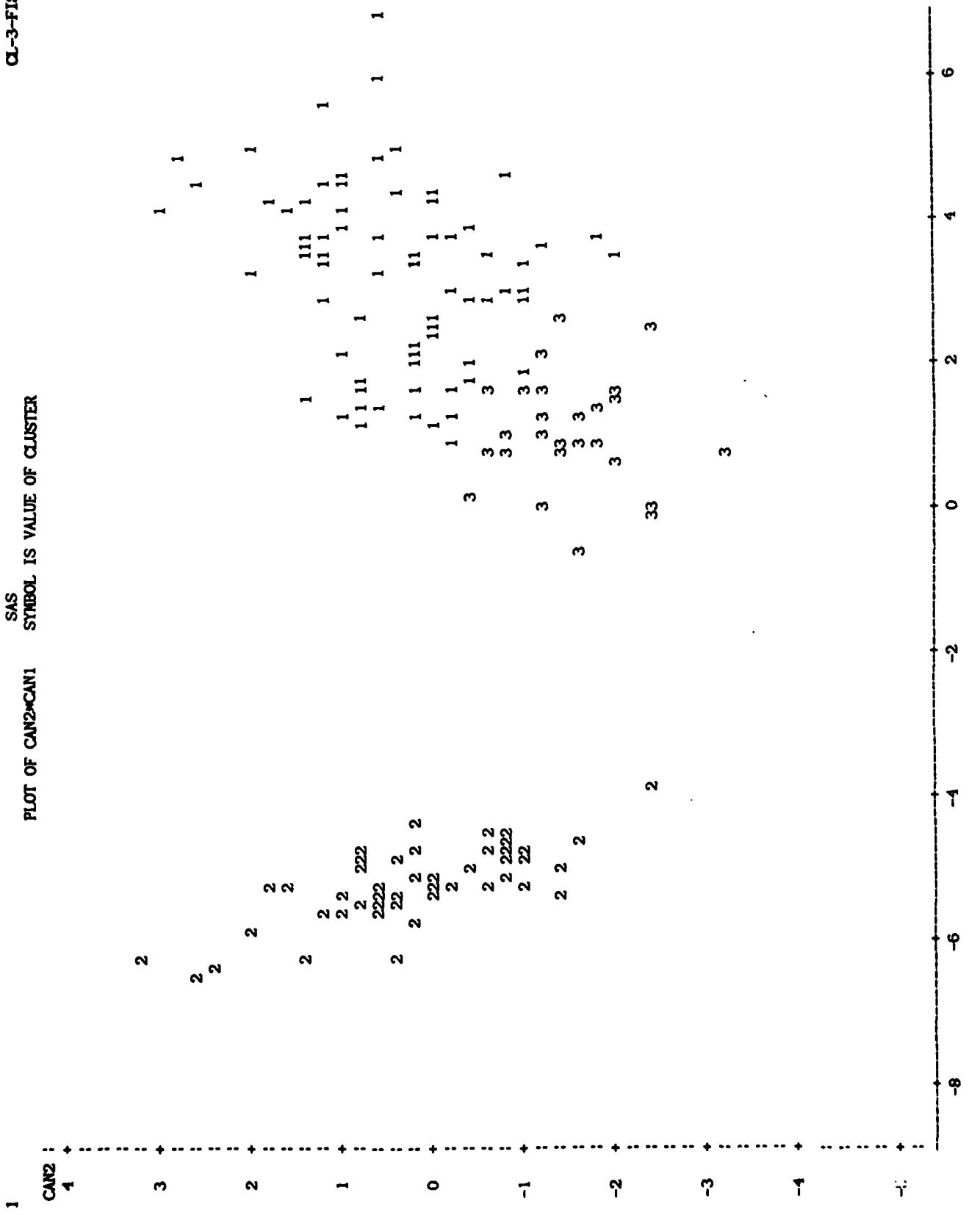
-1579.5

TABLE OF CLUSTER BY GROUP

CLUSTER	GROUP	FREQUENCY			TOTAL
		1	2	3	
1	0	23	49	72	
	0.00	15.33	32.67	48.00	
	0.00	31.94	68.06		
	0.00	46.00	98.00		
2	50	0	0	0	
	33.33	0.00	0.00	33.33	
	100.00	0.00	0.00		
	100.00	0.00	0.00		
3	0	27	1	28	
	0.00	18.00	0.67	18.67	
	0.00	96.43	3.57		
	0.00	54.00	2.00		
TOTAL	50	50	50	150	
	33.33	33.33	33.33	100.00	

CL-3-FISHER

PLOT OF CAN2*CAN1
SAS SYMBOL IS VALUE OF CLUSTER



NOTE: 7 OBS HIDDEN

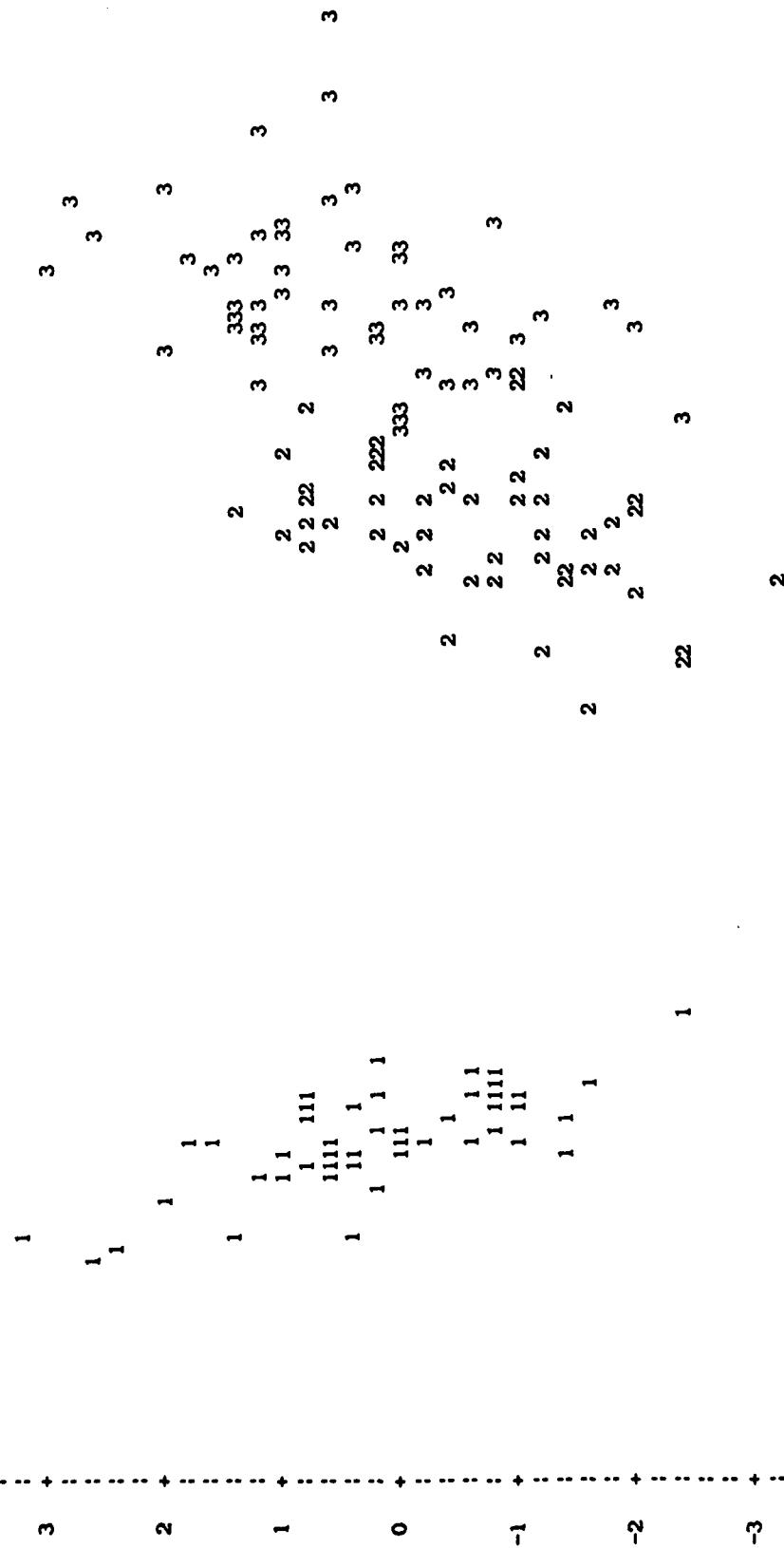
CAN1

43

PLOT OF CAN2*CAN1

SAS
SYMBOL IS VALUE OF GROUP

CL-3-FISHER



NOTE: 7 OBS HIDDEN

CAN1

44

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